Figure 1A

1	CACCAGCAGTAGTAGCAGAAGCGAAGAGCGCAAACGCAACCGCTCTCCCCGCGCGTTGGC	60
61		120
121	. $ \dot{\textbf{ACGCAATTAATGTGAGTTAGCTCACTCATTAGGCACCCCAGGCTTTACACTTTATGCTTC} $	180
181		240
241		300
301		360
361 1	GAGGCAGCAGCTCCACTCAGCCAGTACCCAGATACGCTGGGAACCTTCCCCAGCCAT M	420 1
421 2	GGCTTCCCTGGGGCAGATCCTCTTCTGGAGCATAATTAGCATCATCATTATTCTGGCTGG	480 21
481 22	AGCAATTGCACTCATCATTGGCTTTGGTATTTCAGGGAGACACTCCATCACAGTCACTAC A I A L I I G F G I S G R H S I T V T T	540 41
541 42		600 61
601 62	CATCAAACTTTCTGATATCGTGATACAATGGCTGAAGGAAG	660 81
661 82		720 101
721 102	AGCAGTGTTTGCTGATCAAGTGATAGTTGGCAATGCCTCTTTGCGGCTGAAAAACGTGCAAA VFADQVIVGNASLRLKNVQ	780 121
781 122	ACTCACAGATGCTGGCACCTACAAATGTTATATCATCACTTCTAAAGGCAAGGGGAATGC L T D A G T Y K C Y I I T S K G K G N A	840 141
841 142	TAACCTTGAGTATAAAACTGGAGCCTTCAGCATGCCGGAAGTGAATGTGGACTATAATGC N L E Y K T G A F S M P E V N V D Y N A	900 161
901 162	CAGCTCAGAGACCTTGCGGTGTGAGGCTCCCCGATGGTTCCCCCAGCCCACAGTGGTCTG S S E T L R C E A P R W F P Q P T V V W	960 181
961 182	7 C O V D O O 1 12	1020 201

Figure 1B

1021 202	THE THE THE THE TENT OF THE TE	1080 221
1081 222	THE PROPERTY OF THE PROPERTY O	1140 241
1141 242	The state of the s	1200 261
1201 262		1260 281
1261 282		1320 283
1321		1380
1381	ATCTAGAAGTCTGGAGTGAGCAAACAAGAGCAAAAAAAAA	1440
1441		1500
1501		1560
1561	ATCCCCAGATCTCAGGGACCTCCCCCTGCCTGTCACCTGGGGAGTGAGAGGACAGGATAG	1620
1621		1680
1681		1740
1741		1800
1801		1860
1861		1920
1921		1980
1981	. $. \\$ ATTTCTCAGATGATGTTCATCCGTGAATGGTCCAGGGAAGGACCTTTCACCTTGACTATA	2040
2041	TGGCATTATGTCATCACAAGCTCTGAGGCTTCTCCTTTCCATCCTGCGTGGACAGCTAAG	2100
2101		2160
2161		2220

Figure 1C

2221	CAGTGCTACTACCAACTAGTGGATAAAGGCCAGGGATGCTGCTCAACCTCCTACCATGTA	2280
2281		2340
2341		2400
2401		2460
2461		2520
2521		2580
2581		2640
2641		2700
2701		2760
2761		2820
2821		2880
2881		2940
2941		3000
3001	GAGTCGTATTACAATTCACTGGCCGTCGTTTTACAACGTCGTGACTGGGAAAACCCTGGC	3060
3061	GTTACCCAACTTAATCGCCTTGCAGCACATCCCCCTTTCGCCAGCTGGCGTAATAGCGAA	3120
3121	GAGGCCCGCACCGATCGCCCTTCCCAACAKTTGCGCAGCCTGAATGGCGAATGGCAAATT	3180
3181	GTAAGCGTTAAATTTTGTTAAAATTTTTTTTTTTTTTTT	3240
3241		3300
3301		7

Figure 2

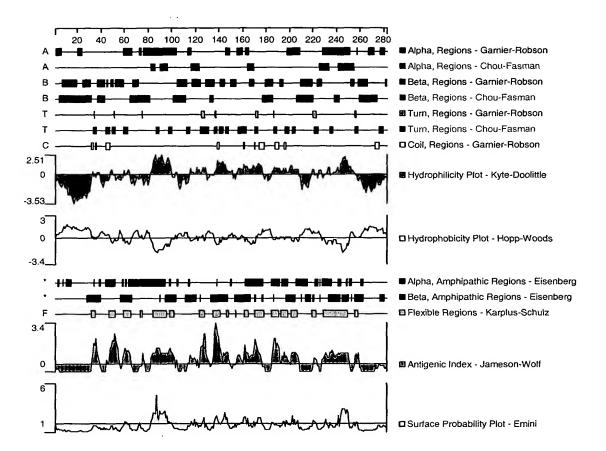


Figure 3A

1	CCACGCGTCCGGAATGAACAACTTTTCTTCTCTTGAATATATCTTAACGCCAAATTTTGA	60
61	GTGCTTTTTTGTTACCCATCCTCATATGTCCCAGCTGGAAAGAATCCTGGGTTGGAGCTA	120
121	. CTGCATGTTGATTGTTTTTTCCTTTTGGCTGTTCATTTTGGTGGCTACTATAAGGA	180
181		240
241 1		300 10
301 11	GAATTGCAGCTTCACCAGATAGCAGCTTTATTCACAGTGACAGTCCCTAAGGAACTGTAC E L Q L H Q I A A L F T V T V P K E L Y	360 30
361 31		420 50
421 51	AACCTTGGAGCAATAACAGCCAGTTTGCAAAAGGTGGAAAATGATACATCCCCACACCGT N L G A I T A S L Q K V E N D T S P H R	480 70
481 71		540 90
541 91	CAAGTCCAAGTGAGGGACGAAGGACAGTACCAATGCATAATCATCTATGGGGTCGCCTGG Q V Q V R D E G Q Y Q C I I I Y G V A W	600 110
501 111	GACTACAAGTACCTGACTCTGAAAGTCAAAGCTTCCTACAGGAAAATAAACACTCACATC D Y K Y L T L K V K A S Y R K I N T H I	660 130
561 L31		720 150
721 .51	GCAGAAGTATCCTGGCCAAACGTCAGCGTTCCTGCCAACACCAGCCACTCCAGGACCCCT A E V S W P N V S V P A N T S H S R T P	780 170
81 71	GAAGGCCTCTACCAGGTCACCAGTGTTCTGCGCCTAAAGCCACCCCCTGGCAGAAACTTC E G L Y Q V T S V L R L K P P P G R N F	840 190
41 91	AGCTGTGTGTCTGGAATACTCACGTGAGGGAACTTACTTTGGCCAGCATTGACCTTCAA S C V F W N T H V R E L T L A S I D L Q	900 210
01 11	AGTCAGATGGAACCCAGGACCCATCCAACTTGGCTGCTTCACATTTTCATCCCCTCCTGC S Q M E P R T H P T W L L H I F I P S C	960 230
61 31		1020 250

Figure 3B

1021 251	TARGET OF THE PROPERTY OF THE	1080 270
1081 271	The state of the s	1140 284
1141		1200
1201		1260
1261	CTGGCCATGAAACTTGCCCCTTCACTGATCTGGACTCACCTCTGGAGCCTATGGCTTTAA	1320
1321	GCAAGCACTACTGCACTTTACAGAATTACCCCACTGGATCCTGGACCCACAGAATTCCTT	1380
1381	CAGGATCCTTCTTGCTGCCAGACTGAAAGCAAAAGGAATTATTTCCCCTCAAGTTTTCTA	1440
1441	AGTGATTTCCAAAAGCAGAGGTGTGTGGAAATTTCCAGTAACAGAAACAGATGGGTTGCC	1500
1501	AATAGAGTTATTTTTTTTATCTATAGCTTCCTCTGGGTACTAGAAGAGGCTATTGAGACTAT	1560
1561	GAGCTCACAGACAGGCTTCGCACAAACTCAAATCATAATTGACATGTTTTATGGATTAC	1620
1621	TGGAATCTTGATAGCATAATGAAGTTGTTCTAATTAACAGAGAGCATTTAAATATACACT	1680
1681	AAGTGCACAAATTGTGGAGTAAAGTCATCAAGCTCTGTTTTTGAGGTCTAAGTCACAAAG	1740
1741	CATTTGTTTTAACCTGTAATGGCACCATGTTTAATGGTGGTTTTTTTT	1800
1801	TTTCCTTTAAAAATTATTGGTTTCTTTTTATTTGTTTTTACCTTAGAAATCAATTATATA	1860
1861	CAGTCAAAAATATTTGATATGCTCATACGTTGTATCTGCAGCAATTTCAGATAAGTAGCT	1920
1921	AAAATGGCCAAAGCCCCAAACTAAGCCTCCTTTTCTGGCCCTCAATATGACTTTAAATTT	1980
1981	GACTTTTCAGTGCCTCAGTTTGCACATCTGTAATACAGCAATGCTAAGTAGTCAAGGCCT	2040
2041	TTGATAATTGGCACTATGGAAATCCTGCAAGATCCCACTACATATGTGTGGAGCAGAAGG	2100
2101	GTAACTCGGCTACAGTAACAGCTTAATTTTGTTAAATTTGTTCTTTATACTGGAGCCATG	2160
2161	AAGCTCAGAGCATTAGCTGACCCTTGAACTATTCAAATGGGCACATTAGCTAGTATAACA	2220
2221	GACTTACATAGGTGGGCCTAAAGCAAGCTCCTTAACTGAGCAAAATTTGGGGCTTATGAG	2280

Figure 3C

2281	AATGAAAGGGTGTGAAATTGACTAACAGACAAATCATACATCTCAGTTTCTCAATTCTCA	2340
2341	TGTAAATCAGAGAATGCCTTTAAAGAATAAAACTCAATTGTTATTCTTCAAAAAAAA	2400
2401	AAAAAA 2406	

Figure 4

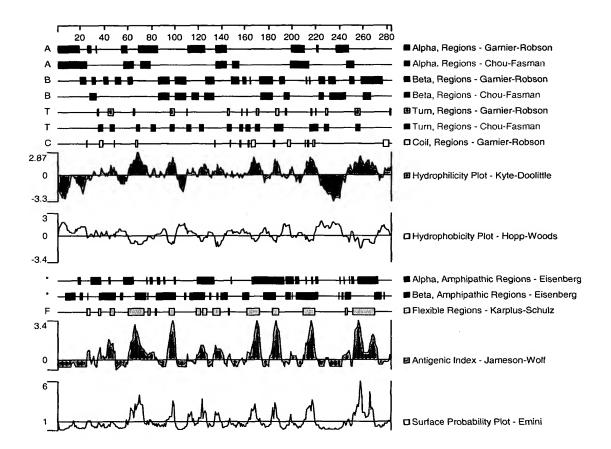


Figure 5A

1 1	G	GCA	CGA	GCT	GTC?	ATC	CGT	FTC	CAT	GCC	GTG.	AGG	TCC	ATT	CAC.	AGA	ACA	CAT	CCA'	TGGCT A	60 2
61 3	C'			TCA(S	GTTI L	rgg: V		rgad S								GGC2 Q		GGC. Q		FGTTI	120 22
121 23	G(G			ACAZ K	AGCC P	TG: V	rcc <i>i</i> Q	AGG(A	CCT: L	rgg [,] V	IGG(G	GGG. E	AGG. D	ACG(A	CAG(A	CAT' F	rct(CCT(C	GTT'	FCCTG L	; 180 42
181 43	T(CTC(P		AGAO T	CAA N	ATG(CAGA E	AGG(CCAT M	rgg: E	AAGT V			TCT' F	TCAC R	G G	GCCI Q	AGT'	rct(s	CTAGC S	240 62
241 63	G1 V		rcc <i>i</i> H		CTA Y	CAC R	EGGA D	CGC G	GAZ K	AGG/ D		AGC(TAT M	rgc <i>i</i> Q	AGAT M	rgcc P	CACA Q	AGT <i>I</i> Y	ATCAA Q	300 82
301 83	GG G	GCA(R	GGAC T	: CAAA K	ACT L	'GG'I V	GAA K	.GGA D	ATTO S	rat: I	TTGC A	CGG <i>I</i> E	G G	GGC(R	GCA1	CTC S	CTCI L	GAC R	GCT L	TGGAA E	360 102
361 103	A.A.	CAT	TAC T	TGT V	GTT L	GGA D		TGG G		CTA Y		GTC C	GCAC R	FGAT I	TTAG S	TTC S	CCA Q	GTC S	TTA Y	CTAC Y	420 122
421 123	CA Q	GA <i>I</i> K	AGGC A	CAT I	CTG W	GGA E	GCT L	aca Q	.GGT V	GTC S	AGC A	ACT L	G G	GCTC S	CAGI V	TCC P	TCT L	'CA'I	TTC S	CATC I	480 142
481 143	GC A	GGG G	ATA Y	TGT V	TGA D	TAG R	AGA D		CCA Q	GCT L		CTG C		GTC S	CTC S	GGG G	CTG W	GTT F	CCC P	CCGG R	540 162
541 163	CC P	CAC T	AGC A	GAA K	GTG W	GAA K	AGG G	TCC P	ACA Q	AGG G	ACA Q	GGA D	TTT L	GTC S	CAC T	AGA D	CTC S	CAG R	GAC T	AAAC N	600 182
601 183	AG R	AGA D	CAT M	GCA H	TGG(G	CCT L	GTT' F	TGA D	TGT V	GGA E	GAT I	CTC S	TCT L	GAC T	CGT V	CCA Q	AGA E	GAA N	CGC A	CGGG G	660 202
661 203	AG S	CAT I	ATC S	CTG' C	TTC(S	CAT(M	GCG(R	GCA' H	TGC' A	TCA H	TCT(L	GAG S	CCG R	AGA E	GGT V	GGA. E	ATC S	CAG R	GGT. V	ACAG Q	720 222
721 223	AT.	AGG G	AGA(CTG: W	GAG <i>I</i> R	AAG/ R	AAAG K	GCA(CGG; G	ACA Q	GGC A	AGG G	TAA K	AAG. R	AAA. K	ATA' Y	TTCC S	CTC S	TTC: S	ACAC H	780 242
781 243	AT'	TTA Y	TGA(CTC(S	CTTI F	P P	AAG7 S	rct(L	CTC(S	GTT' F	- FATO M	GGA' D	TTT' F			CCT(L			CGT(V		840 262
841 263	CC(CTG C	CAG <i>I</i> R	AGCO A	CAAC K	CT: L	rgtc V	SATO M	GGGZ G	AAC' T	FCT(L	GAA K	ATT(L	GCA(GAT'	rct(GGGG	GA(E	GT(V	GCAT H	900 282
901 283	TTT F	rgt: V	AGAC E	GAAG K	GCCC P	CAT H	TAGC S	CTI L	rcti L	rcac Q	GATO	CTC:	rggz G	AGG(GTC(CACA T		L L	CAA/ K	AAAG K	960 302

Figure 5B

961 303	TARGOLD TO THE PROPERTY OF THE	1020 319
1021	TGCCTGCTCTCTGCTTTCAGAATTGAGAGACGCCCGGAAACACGCAGGTACCAA	1080
1081		1140
1141		1200
1201		1260
1261		1320
1321	GTGCAAAGGGGAACGAAGTGAAGTTAGCAGGAACTGGTGGGTG	1380
1381	GGAGTCACTCAAGGTCTCACAAAGTCAAATAGAGGGCTTACGTGGGAGGGCAGTGGTAGG	1440
1441	GCTGGGTGAACATCTCATGGTTGAGCATCTCCAAGCATCAGTGAGGCACGGGGGCTGCCC	1500
1501	TGGAGAAGGTACATGGCTGGGATAGTGGGACTGGCCGGATCCTACCCGGAGCCAGTC	1560
1561	TGCAGTGGGAGGGTCGACCTCTTGCTCCAGCCCAGATTTCGTCTTCAGTAACTCATGCTT	1620
1621	CCTCTCTCCCCCACCGCACCCCAGTGGAGGTGACTCTGGATCCAGAGACGGCTCACCCGA	1680
1681	AGCTCTGCGTTTCTGATCTGAAAACTGTAACCCATAGAAAAGCTCCTCAGGAGGTGCCTC	1740
1741	ACTCTGAGAAGAGTTTACAAGGAAGAGTGTGGTGGCTTCTCAGGGTTTCCAAGCAGGGA	1800
1801	AACATTACTGGGAGGTGGACGTGGGACAAAATGTAGGGTGGTATGTGGGAGTGTCTCGGG	1860
1861		1920
1921	TCAGACTGACAACAGAACATTTGTATTTCACATTCAATCCCCATTTTATCAGCCTCCCCC	1980
1981	CCAGCACCCCTCCTACACGAGTAGGGGTCTTCCTGGACTATGAGGGTGGGACCATCTCCT	2040
2041		2100
2101		2160
2161		2220

Figure 5C

2221	GACCCAGACACAGCCAAGGGAGAGTGCTCCCGACAGGTGGCCCCAGCTTCCTCCGGAG	2280
2281		2340
2341	GAGCCCTGCAGCAGCGGCAGTCACAGCTTCCAGATGAGGGGGGGATTGGCCTGACCCTGTG	2400
2401		2460
2461	TTGTGAAAACTCCATCCAGCTAAGCGATCTTGAACAAGTCACAACCTCCCAGGCTCCTCA	2520
2521	TTTGCTAGTCACGGACAGTGATTCCTGCCTCACAGGTGAAGATTAAAGAGACAACGAATG	2580
2581	TGAATCATGCTTGCAGGTTTTGAGGGCCACAGTGTTTTGCTAATGGATGTGTTTTTATGATT	2640
2641	ATACATTTTCCCCACCATAAAACTCTGTTTGCCTTAATTCCCACATTAATTTAACTTTTC	2700
2701	CTCCTATACCCAAATCCACCCATGGAATAGTTAATTGGAACACCTGCCTTTGTGAGGCTC	2760
2761	CAAAGAATAAAGAGGAGGTAGGATTTTTCACTGATTCTATAAGCCCAGCATTACCTGATA	2820
2821		2880
2881	TTAACACAGACACAAAATTCTAAATAAAATTTTAACAAATTAAACTAAACAATATATTT	2940
2941	AAAGATGATATAACTACTCAGTGTGGTTTGTCCCACAAATGCAGAGTTGGTTTAATAT	3000
3001	TTAAATATCAACCAGTGTAATTCAGCACATTAATAAAGTAAAAAAAA)59

Figure 6

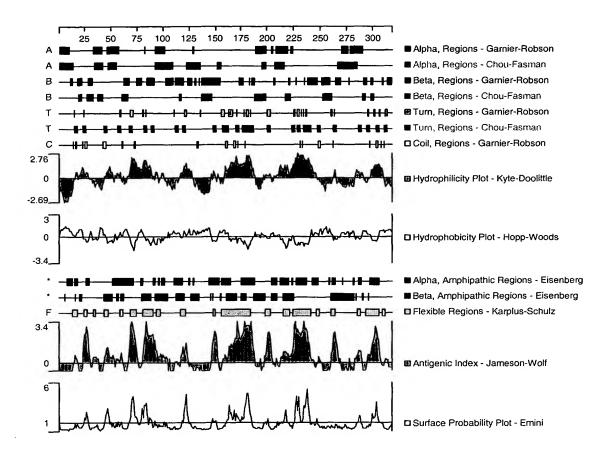


Figure 7A

1	NNCACGAGCCTGTGCCCCTGGAAAGGTTGGAGACTTGGGGGACGACTGGAGAATTGCCAT													
61		120												
121 1	TGCTCTGGGTGCTCATGGAACCAGCTGCTGCCCTGCACTTCTCCCGGCCAGCCTCCCTC	180 16												
181 17	TCCTCCTCAGCCTGTGTGCACTGGTCTCAGCCCAGTTTACTGTCGTGGGGCCAGCTA L L L S L C A L V S A Q F T V V G P A N	240 36												
241 37	ATCCCATCCTGGCCATGGTGGGAGAAACACTACATTACGCTGCCATCTGTCACCCGAGA PILAMVGENTTLRCHLSPEK	300 56												
301 57	AAAATGCTGAGGACATGGAGGTGCGGTGGTTCCGGTCTCAGTTCTCCCCCGCAGTGTTTG N A E D M E V R W F R S Q F S P A V F V	360 76												
361 77	TGTATAAGGGTGGGAGAGAACAGAGGAGCAGATGGAGGAGTACCGGGGAAGAATCA Y K G G R E R T E E Q M E E Y R G R I T	420 96												
421 97		480 116												
481 117	CCCAGGAGAATGGGATCTACCGCTGTTACTTCCAAGAAGGCAGGTCCTACGATGAGGCCAQENGENGENGENGENGENGENGENGENGENGENGENGENGE	540 136												
541 137	TCCTACGCCTCGTGGTGGCAGGCCTTGGGTCTAAGCCCCTCATTGAAATCAAGGCCCAAG L R L V V A G L G S K P L I E I K A Q E	600 156												
601 157	AGGATGGGAGCATCTGGAGGTGCATATCTGGAGGGTGGTACCCAGAGCCCCTCACAG DGSIWLECISGGWYPEPLTV	660 176												
661 177	TGTGGAGGGACCCCTACGGTGAGGTTGTGCCCGCCCTGAAGGAGGTTTCCATCGCTGATG W R D P Y G E V V P A L K E V S I A D A	720 196												
721 197	CTGACGGCCTCTTCATGGTCACCACAGCTGTGATCATCAGAGACAAGTATGTGAGGAATG D G L F M V T T A V I I R D K Y V R N V	780 216												
781 217	TGTCCTGCTCTGTCAACAACACCCTGCTCGGCCAGGAGAAGGAAACTGTCATTTTTATTC S C S V N N T L L G Q E K E T V I F I P	840 236												
341 237	CAGAATCCTTTATGCCCAGCGCATCTCCCTGGATGGTGGCCCTAGCTGTCATCCTGACCG E S F M P S A S P W M V A L A V I L T A	900 256												
901 257		960 276												

Figure 7B

961 277	TCAGCATCTGTTGCATCAAGAAACTTCAAAGGGGAAAAAAAA	1020 296
1021 297	AAGTTGAACAAGAGAAAAGAAATTGCACAGCAACTTCAAGAAGAATTGCGATGGAGAA VEOEEKEIA OOLOEELRWRR	1080
	· - 2 1 - 1 - 1 - 2 - 2 - 2 - 1 - 1 -	316
1081 317	GAACATTCTTACATGCTGCTGATGTGGTCCTGGATCCAGACACCGCTCATCCCGAGCTCT T F L H A A D V V L D P D T A H P E L F	1140 336
1141 337	TCCTGTCAGAGGACCGGAGAAGTGTGAGGCGGGGCCCCTACAGGCAGAGAGTGCCTGACA L S E D R R S V R R G P Y R Q R V P D N	1200 356
1201 357	ACCCAGAGAGATTCGACAGTCAGCCTTGTGTCCTGGGATGGGAGAGCTTCGCCTCAGGGA PERFDSQPCVLGWESFASGK	1260 376
1261 377	AACATTACAGGGGAAACTTCACAGAGTGGGGACCCACCAGAGCCTATAGAATCAATTCCT H Y R G N F T E W G P T R A Y R I N S L	1320 396
1321 397	TGGACTCACAGCCATGCAGAAAGCCCTGGCCATCTCAGCAGCCACCGCACAACCCCCCTA D S Q P C R K P W P S Q Q P P H N P P N	1380 416
1381 417	ATGAAAGACACGCCCTCCCCCTCTGGTCACGTAAGAGAACATCTTCCAGCTGCCTTTT ERHALLPSGHVREHLPAAFF	1440 436
1441 437		1500 455
1501		1560
1561		1620
1621		1680
1681		1740
1741		1800
1801		1860
1861	GTCCTCCCTGAGAGGATTCTCCCTTTGAAGGAGTCCCTTTGCCGGGTGGGCGTCTTCCT	1920
1921		1980
1981		2040

Figure 7C

2041	CAGCCCCATCTTCATCTGCCCTGCACTCACAGGAGCCAGTGGGGTCATGGTGCCTGAAGA	2100
2101		2160
2161		2220
2221		2280
2281		2340
2341		2400
2401		2460
2461		2520
2521		2580
2581		2640
2641		

Figure 8

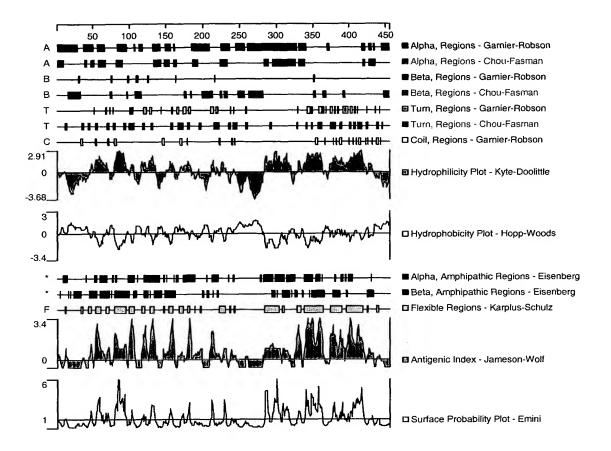


Figure 9A

1	C	GAT'	TCG	GCT(CCAZ	AAC	rcco	GCC	CTC	GCAC	GCCG	ATC	GGA	CTC	TGG	GCC	GCG	GTO	GGC	CACCG	60
61	C	GCG(CAG	CTAC	GGG <i>I</i>	AGCC	CGAC	GAAC	CGC	CGGC	CGAC	GCC	CGA	.GGA	rcgc	CCA	GAG	CGC	'GAG	GGTC	120
121	G	CTG	CGC	CTCC	GCAC	GAGC	CCGG	GAGC	:CG <i>I</i>	AGTO	GAG	CCG	GGC	GCC	:CGG	GCT	GCC	TGG	AGA	CGCC	180
181 1	G7	rga(CTT	rga <i>z</i>	AGTO	STAI	ACTI	CAA	GAC	CAGA	ATGG	GCG	CAT M	GCG R	GGA E	GAT I	CGT V	GTG W	GTA Y	CCGG R	240 8
241 9	GT V	GAO T		ATGG G	G G	GCAC T	CAT I	CAA K	.GCA Q	AAA K	GAT I	CTT F	CAC T	CTT F	CGA D	CGC A	CAT M	GTT F	CTC S	CACC T	300 28
301 29	A <i>P</i> N	CT? Y	CTC S	ACA H	CAT M	'GGA E	GAA N	CTA Y	.CCG R	CAA K	GCG R	AGA E	GGA D	CCT L	GGT V	GTA Y	CCA Q	GTC S	CAC T	TGTG V	360 48
361 49	AG R	GC'I L	GCC P	CGA E	GGT V	'CCG R	GAT I	CTC S	AGA D	.CAA N	TGG G	TCC P	CTA Y	TGA E	GTG(CCA' H	TGT V	GGG G	CAT I	CTAC Y	420 68
421 69	GA D	CCG R	CGC A	CAC T	CAG R	GGA E	GAA K	GGT V	GGT V	CCT L	GGC. A	ATC. S	AGG(G	CAA N	CATO	CTT(F	CCT(L	CAA N	CGT V	CATG M	480 88
481 89	GC A	TCC P	TCC P	CAC T	CTC S	CAT I	TGA E	AGT V	GGT V	GGC A	TGC' A	TGA D	CAC:	ACC. P	AGC(A	P	CTT(CAG S	CCG(R	CTAC Y	540 108
541 109	CA Q	AGC A	CCA Q	GAA N	CTT F	CAC T	GCT L	GGT V		CAT I	CGT(V	GTC' S	rgg: G	AGG2 G	AAA/ K	ACC) P	AGCI A	ACC(P	CATO M	GGTT V	600 128
601 129	TA Y	TTT F	CAA K	ACG R	AGA D	TGG G	GGA. E	ACC P	AAT(CGA(CGC A	AGT(V	GCC(CCTA L	ATC <i>I</i> S	AGA(E	GCC <i>I</i> P	ACC? P	AGC! A	IGCG A	660 148
661 149	AG S	CTC S	CGG G	CCC P	CCT. L	ACA(Q	GGA	CAG(S	CAGO R	GCC(CTT(F	CCG(R	CAG(S	CCTI L	rcte L	GCAC H	CCGT R	rga(D	CCT(GGAT D	720 168
721 169	GA D	CAC T	CAA K	GATO M	GCA(Q	GAA(K	GTC S	ACTO L	STC(CCT(L	CCTC L	GGA(D	GCC A	EGAC E	SAAC N	CGG R	GGT	rgg(G	GCG <i>I</i> R	ACCC P	780 188
781 189	TAC Y	CAC T	GGA(GCG(R	P P	CTC(S	CCG!	rgg(G	CCT(L	GAC(CCC <i>P</i> P	AGAT D	P	CAAC N	CATC	CTC L	CTC L	CAC Q	GCC <i>P</i> P	AACC T	840 208
841 209				CATA I		AGA(E	GACO T	GTC V	GTC V	GAGO S	CCGI R	GAG E	TTT F	PCCC	CGC R	TGG W	GTC V	CAC H	AGC S	EGCC A	900 228
901 229	GA0 E	GCC(CAC(T	CTAC Y	CTTC F	CCT(GCGC R	CCAC H	:AGC S	CCGC R	ACC T	CCG P	AGC S	AGT S	GAC D	GGC G	ACT T	GTG V	GAA E	GTA V	960 248
961 249	CGT R	rgco A	CCT(L	GCTC L	CACC T	CTGC W	GACC T	CTC L	AAC N	CCA P	.CAG Q	ATC I	GAC D	AAC N	GAG E	GCC A	CTC L	TTC F	AGC S	TGC C	1020 268

Figure 9B

1021 269	GAGGTCAAGCACCCAGCTCTGTCGATGCCCATGCAGGCAG	1080 288
		200
1081 289	AAAGGACCCAAAATTGTGATGACGCCCAGCAGAGCCCGGGTAGGGACACAGTGAGGATT K G P K I V M T P S R A R V G D T V R I	1140 308
1141		1200
309	L V H G F Q N E V F P E P M F T W T R V	328
1201 329		1260
		348
1261 349	GTTCCCGCCGAGCTCAATGGCTCCATGTATCGCTGCACCGCCCAGAACCCACTGGGCTCC V P A E L N G S M Y R C T A Q N P L G S	1320 368
1321	ACCGACACGCACACCCGGCTCATCGTGTTTGAAAACCCAAATATCCCAAGAGGAACGGAG	1380
369	TDTHTRLIVFENPNIPRGTE	388
1381 389	GACTCTAATGGTTCCATTGGCCCCACTGGTGCCCGGCTCACCTTGGTGCTCGCCCTGACA D S N G S I G P T G A R L T L V L A L T	1440 408
1 4 4 7		100
409	GTGATTCTGGAGCTGACGTGAAGGCACCCGCCCCGGCCACTCCATCAGGCACTGACATCT V I L E L T *	1500 415
1501		1560
1561	ATCTGTGTCTTGGCTTCTTCAGTCGGTTTAATTAAAACAAAC	1620
1601		
1621	AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	1680
1681	АААААААААААААААААААААААААААААААА	

Figure 10

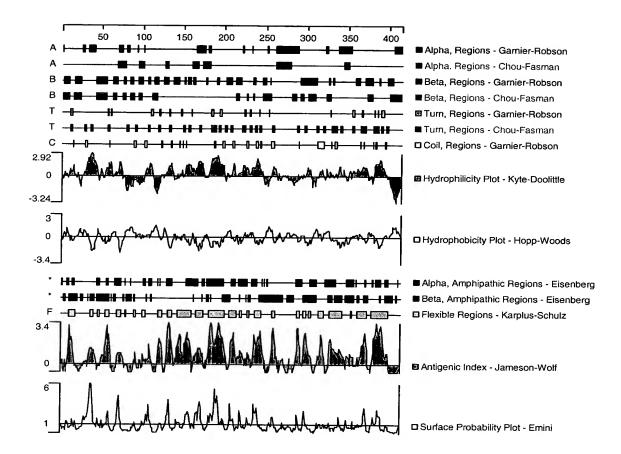


Figure 11

1	CACGAGCCTGTGCCCCTGGAAAGGTTGGAGACTTGGGGGACGACTGGAGAATTGCCATTT	60
61	GAGGACCAAAGGAGAAAAGAAACTACACGCTAATTCTAGAAGGCCTCCTGTCCCTG	120
121 1	The state of the s	180 16
181 17	The state of the s	240 36
241 37	A A CONTROL OF THE CO	300 56
301 57	The state of the s	360 76
361 77	TO THE CONTROL OF THE	420 96
421 97	TTTGTGAGCAAAGACAGCAGGGGCAGCGTGGCCCTGATCATACACAATGTCACAGCCGAG F V S K D S R G S V A L I I H N V T A E	480 116
481 117	GATAACGGCATCTACCAGTGTTACTTCCAAGAAGGCAGGTCCTGCAATGAGGCCATCCTG D N G I Y Q C Y F Q E G R S C N E A I L	540 136
541 137		600 156
601 157		660 160
661		720
721		780
781		840
841		900
901		960
961	AAGTAGAGCGTTTTATTAAAGCAAGACTTAATACAGAAGCAAAAAAAA	.9

Figure 12

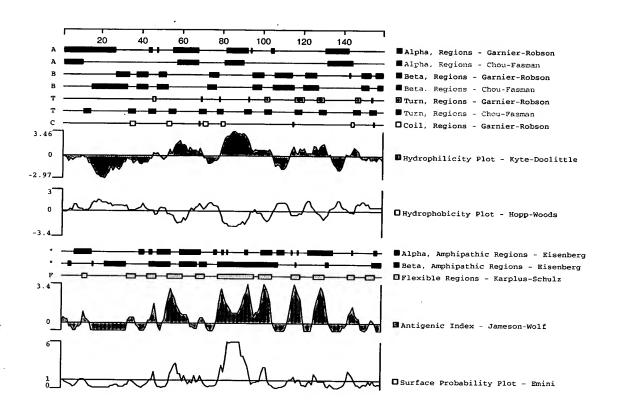


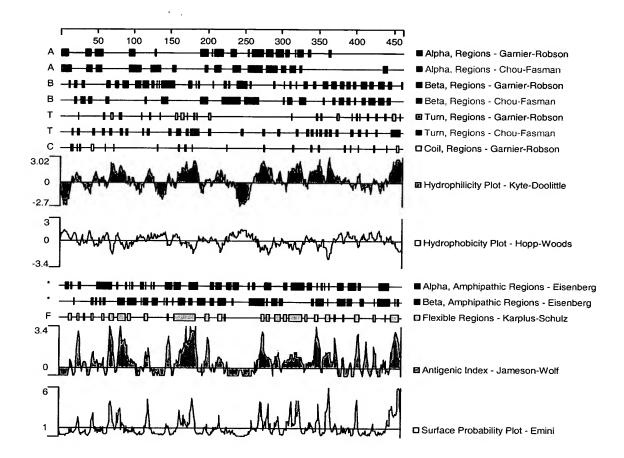
Figure 13A

1 1	The state of the s	60 18
61 19	The state of the s	120
121 39	The state of the s	180 58
181 59	CAGTTCTCTAGCGTGGTCCACCTCTACAGGGACGGGAAGGACCAGCCATTTATGCAGATGQ F S S V V H L Y R D G K D Q P F M Q M	240 78
241 79		300 98
301 99		360 118
361 119	CAGTCTTACTACCAGAAGGCCATCTGGGAGCTACAGGTGTCAGCACTGGGCTCAGTTCCTQSYYQKAIWELQVSALGSVP	420 138
421 139		480 158
481 159	TGGTTCCCCCGGCCCACAGCGAAGTGGAAAGGTCCACAAGGACAGGATTTGTCCACAGAC W F P R P T A K W K G P Q G Q D L S T D	540 178
541 179	TCCAGGACAACAGAGACATGCATGGCCTGTTTGATGTGGAGATCTCTCTGACCGTCCAAS R T N R D M H G L F D V E I S L T V Q	600 198
601 199	GAGAACGCCGGGAGCATATCCTGTTCCATGCGGCATGCTCATCTGAGCCGAGAGGTGGAA E N A G S I S C S M R H A H L S R E V E	660 218
661 219		720 238
721 239	GTACTGGGAATACTCTGCTGTGGCCTATTTTTTGGCATTGTTGGACTGAAGATTTTCTTC V L G I L C C G L F F G I V G L K I F F	780 258
781 259	TCCAAATTCCAGTGGAAAATCCAGGCGGAACTGGACTGG	840 278
841 279	GAATTGAGAGACGCCCGGAAACACGCAGTGGAGGTGACTCTGGATCCAGAGACGGCTCAC E L R D A R K H A V E V T L D P E T A H	900 298
901 299		960 318

Figure 13B

961	CC	TCA	CTC	TGA	GAA	GAG	ATI	'TAC	AAG	GAA	GAG	TGI	GGT	GGC	ттс	TCA	GAG	ттт	'CCA	AGCA	1020
319	P	H	S	E	K	R	F	T	R	K	S	V	V	A	S	Q	S	F	Q	A	338
1021	GG	GAA	ACA	ጥጥል	СТС	:CCA	ССТ	CCA	CGG	acc.	aca	~ n n	. ת א יחי		'CMC	יכככ		ICCC	3 C C	GTGC	1000
339	G	K	H	Y	W	E	V	D	G	G	H H	N.CAA	K	R	W	R	V V	G G	V V	C	1080 358
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1081	CG	CCA	тса	тст	CCA	C ል C	GAG	CVV	CCA	ርሞል	~~~	C 2 C	mmm.	·	mco	CC 2		maa	am s	CTGG	1140
359	R	D	D	V	D	R	R	K	E	Y	V	T	L	S	P	D	H	G	GTA Y	W	1140 378
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1141	СT	ירכייי	CAG	አርጥ	~ A A	חרר	7 C 7	7 C 7	mmm	OM N	mmm	a 2 a	3 mm		.					CGTC	
379	v	L	R	L	N N	G	E E	H	L	GIA Y	F	T T	ATT L	AAA N	P	CCG R	TTT F	TAT	CAG S	V	1200 398
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1201	ጥጥ	רככ	CAG	Gac	רככ	אככי	ጥአሮ	א א א	ת א	አርር		C mm	ccm			mo a			~~	CATC	1050
399	F	P	R	T	P	P	T.	K	I	G G	V	F	L	GGA D	Y Y	TGA	C	TGG G	GAC T	I	1260 418
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1261	ጥር	_ 	СФФ	• ግአአ	~ አጥ	יגגג	TC N	~~~	ome.	-cmr	nam	י אינות אינו	m » <		~ ~ ~	.				AGGC	
419	S	F	F	N N	I	AAA N	D D	O	S	L L	IAT I	Y	TAC T	CCT L	GAC. T	ATG' C	TCG R	GTT F	TGA. E	AGGC G	1320 438
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1321	ጥጥ	א היהי	C N C (י מיחיר	יים אי	nca	י מיחיר	TICC(2ma	•		mc 3 .								
439	L	L	R	P	Y	I	E	Y	P	S	Y Y	N N	TGA(GCA. O	AAA' N	TGG/ G	AAC' T	P P	CAGA R	AGAC D	1380 458
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1381	7 7 6	CC3:	A (" A (л <i>с</i> тт	7.CM/		73 C				200	~~~								
459		O	O	*	HGT.	CT	CT	JACZ	AGG(JAAC	CAC	JGC	CC'I"	rcc.	rcc	CCA	3GG(GTG2	AAA:	rgta	1440 462
		-	-																		402
1441	CC	ለ ጥር ን	እ አ <i>ጥ</i> ረ	•	י שכינ	7070	• • • • • • • • • • • • • • • • • • • •	namr	n como												
1441	GGZ	A I G	MAI (AC	1100	CAC	.A1	ICT.	rC1"	LTAC	الم	ATA.	I.I.Y	AGG".	rcro	J'I'C'.	rccc	CAGA	ATCO	CAAA	1500
1501	GT	CCC	GCAC	CAC	3CC(3GCC	CAAC	GTC	GC1	rTCC	AGA	ATG	AAGO	GGG	GAC:	rggc	CTC	STCC	CACA	ATGG	1560
1561	GAG	GTC!	\GG1	GTC	CATO	GC7	GCC	CTC	AGC	TGG	GAG	GGI	\AG <i>I</i>	AAGO	CTC	SACA	TTI	CAI	TTT	GTT	1620
1621	TGO	CTCI	CAC	TCC	ATC	TGG	· CTA	AGT	GAT	· CTT	'GAA	ATA	ACCA	ACCI	стс	AGG	· TGA	AGZ	ACC	GTC	1680
																				.010	1000
1681	AGC	רעענ	יייירי	יראיי	ירידר	מרמי	.ccc	ייייכייי	ССТ	עיייטי	Слп	א מוחי	Cm2		יאאר	10777			m = =	TGC	1540
	1.00	14 1 I		CAL		C.F.		.191	. 331	GIA	CMI	IAA	1G 17	JAU	AAC	GAA	1161	GAA	TAA	TGC	1740
1741	m	. ~ -																			
1741	TTF	AGAT	.CTT	TT:A:	GAI	GAC	AGA	GTG	TAT	'CCT	TAA	'GG'I	TTC:	TTC	TTA:	'ATA	TTA	CAC	TTT	CAG	1800
1801	יד א מידי	2 2 2	מממ	222	א א א	א א א	7 7 7	7 7 7	7 7 7	222	7 7 7	1	022								

Figure 14



Human B cell proliferation assay

Figure 15

